

WEST Search History

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DATE: Monday, March 01, 2004

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		<i>DB=DWPI; THES=ASSIGNEE; PLUR=YES; OP=OR</i>	
<input type="checkbox"/>	L5	19980918)	0
		<i>DB=USPT; THES=ASSIGNEE; PLUR=YES; OP=OR</i>	
<input type="checkbox"/>	L4	19980918)	0
		<i>DB=PGPB,USPT,DWPI; THES=ASSIGNEE; PLUR=YES; OP=OR</i>	
<input type="checkbox"/>	L3	L2 not genentech.asn.	2
<input type="checkbox"/>	L2	19980918)	367
<input type="checkbox"/>	L1	PRO1157 or DAP10 or Zsig16	542

END OF SEARCH HISTORY

US-09-127-946-7
; Sequence 7, Application US/09127946
; Patent No. 6416973
; GENERAL INFORMATION:
; APPLICANT: Bakker, Alexander B.H.
; APPLICANT: Phillips, Joseph H.
; APPLICANT: Lanier, Lewis L.
; TITLE OF INVENTION: Mammalian Cell Membrane Proteins;
; TITLE OF INVENTION: Related Reagents
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: DNAX Research Institute
; STREET: 901 California Avenue
; CITY: Palo Alto
; STATE: California
; COUNTRY: USA
; ZIP: 94304-1104
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/127,946
; FILING DATE: 31-JUL-1998
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/089,168
; FILING DATE: 12-JUN-1998
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/069,692
; FILING DATE: 16-DEC-1997
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/069,639
; FILING DATE: 15-DEC-1997
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/063,717
; FILING DATE: 29-OCT-1997
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/054,430
; FILING DATE: 01-AUG-1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Ching, Edwin P.
; REGISTRATION NUMBER: 34,090
; REFERENCE/DOCKET NUMBER: DX0763X
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650)852-9196
; TELEFAX: (650)496-1200
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 451 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 63..338
; FEATURE:
; NAME/KEY: mat_peptide
; LOCATION: 117..338
JS-09-127-946-7

Query Match 96.3%; Score 416.8; DB 4; Length 451;
Best Local Similarity 99.3%; Pred. No. 8.2e-114;
Matches 429; Conservative 0; Mismatches 2; Indels 1; Gaps 1;

Qy	2	CTCTGGACCACAGTCCTCTGCCAGACCCCTGCCAGACCCAGTCCACCATGATCCATCTG	61
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Qy	62	GGTCACATCCTCTTCTGCTTTTGTCTCCAGTGGCTGCAGCTCAGACGACTCCAGGAGAG	121
Db	75	GGTCACATCCTCTTCTGCTTTTGTCTCCAGTGGCTGCAGCTCAGACGACTCCAGGAGAG	134
Qy	122	AGATCATCACTCCCTGCCTTTTACCCTGGCACTTCAGGCTCTTGTTCGGATGTGGGTCC	181
Db	135	AGATCATCACTCCCTGCCTTTTACCCTGGCACTTCAGGCTCTTGTTCGGATGTGGGTCC	194
Qy	182	CTCTCTCTGCCGCTCCTGGCAGGCCTCGTGGCTGCTGATGCGGTGGCATCGCTGCTCATC	241
Db	195	CTCTCTCTGCCGCTCCTGGCAGGCCTCGTGGCTGCTGATGCGGTGGCATCGCTGCTCATC	254
Qy	242	GTGGGGGCGGTGTTCTGTGCGCACGCCCCAGCCCGAGCCCCGCCCCAAGATGGCAAAGTC	301
Db	255	GTGGGGGCGGTGTTCTGTGCGCACGCCCCAGCCCGAGCCCCGCCCCAAGATGGCAAAGTC	314
Qy	302	TACATCAACATGCCAGGCAGGGGCTGACCCCTCCTGCAGCTTGGACCTTTGACTTCTGACC	361
Db	315	TACATCAACATGCCAGGCAGGGGCTGACCCCTCCTGCAGCTTGGACCTTTGACTTCTGACC	374
Qy	362	CTCTCATCCTGGATGGTGTGTGGTGGCACAGGAACCCCGCCCCAACTTTTGGATTGTAA	421
Db	375	CTCTCATCCTGGATGGTGTGTGGT-GCACAGGAACCCCGCCCCAACTTTTGGATTGTAA	433
Qy	422	TAAAACAATTGA	433
Db	434	TAAAACATTTGA	445